



>SEQID#1_SGP037

ATGTCAACAGCTGCCCTTAATTACTTTGGTCAGAAAGTGGTGGGAACCCAGGTGAGAGAGTGCTGCTAAGCTCCGCGCTGCTGCAGGACGACA
GGCGGGTGACACCCACGTCGCCACAGCTCCACTTCAGAGCCTAGGTGTCTCGGTTTGACCCAGATGGTAGTGGGAGTCCAGCTACCTGGGACAA
TTTGGGATCTGGGATAACCGCATTTGATGAGCCAAATCTGCTGCCACCCAGCAATTAGTATGGCAAGCCAAATGCCCAAATCAGCTTGGAAATG
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GCAACAGTAGCCCTATTGCGAGATGGTATTGAACCTGTTGTAGCCAGTGTGGGACAGCCGGGCTATTTTGTGTAGAAAGGAAACCCCATGA
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TAAATTACATCATGCTGATGACAGCTTCCCTGGTCCCTCACACAGATGGAAATTAACCTTCATGGTGAATAGTCAAGAGATTTGTGACTTTGTCAATC
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TGGTGCCCTGGGGAATAATAAGAACTCTGAAATCAACTTCTCATTCAGCAGAAAGCTTTGCCCTCCAGTGGACGATGGGCGCTGA

Fig. 1

>SEQID#2_SGP037
MSTAALITLVRSGGNQVRRRVLLSSRLQDDRRVPTICHSTSEPRCSRFPDGGSPATWDFGIWDFNDRIDEPILLPSIKYKPIPKISLENVGCASQIG
KRKENEDRFDFAQLTDEVLYTAVYDGHGCPAAADFCHTHMEKCTMDLLPKEKNLETLTLAFLHDKAFSSHARLSADATLLTSGTTATVALLRDGIEL
VVASVGDRAILCRKGKPMKLTIDHTPERKDEKERKKCGGFVAVNSLGQPHVNGRLAMTRSIGDLDLKTSGVIAEPETKRKLHHADDSEFLVLTIDGI
NFMVNSQEICDFVNCCHDPNEAAHA VTEQAIQYGTEDNSTAVVVFPGAWGKYKNSEINFSSRSFASGRWA

Fig. 2